Reply to Office action of September 3, 2008

Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- (Currently amended) A method of performing <u>two-dimensional</u> Nuclear Magnetic Resonance (NMR) spectroscopy on a hyperpolarized sample, which method comprises the steps of:
 - hyperpolarizing a sample which comprises a first nuclear species (I) and a second nuclear species (S), with the Hamiltonian $H = H_S + H_{IS} + H_{I}$ using DNP, wherein the NMR active nuclei receive hyperpolarization and transformation of the sample to a liquid state;
 - performing two-dimensional NMR spectroscopy on the sample and thereby producing at least two NMR spectra with the use of a sequence of rf-pulses, wherein the pulse pulse sequence comprises at least two rf-pulses, either on the same nuclei or on different nuclei, and wherein the pulse pulse sequence is adapted for a hyperpolarized sample in such a way that it uses a single scan, an efficient trajectory in a t_S-t_{IS} plane and produces a square array of observed points in a square portion of a two time space, thereby producing at least two NMR spectra;
 - analysing the at least two of the NMR spectra in order to obtain a characterization of the sample. For to obtain an interim result to be used in the NMR spectroscopy step
- 2. (Cancelled)
- 3. (Cancelled)
- 4. (Cancelled))
- 5. (Cancelled)
- 6. (Cancelled)
- 7. (Cancelled)

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- 8. (Cancelled)
- 9. (Cancelled)
- 10. (Cancelled)
- 11. (Cancelled)
- 12. (Cancelled)
- 13. (Cancelled)
- 14. (Currently amended) The NMR spectroscopy method according to claim <u>18</u>, wherein the pulse sequence <u>spans a trajectory in a two-dimensional evolution time space (t_S, t_{IS}), said in performed on a hyperpolarized sample which comprises a first nuclear spin species (I) and a second nuclear spin species (S), the pulse sequence comprises the step of:</u>
 - (300) starting from the point (0,0), with an 90° pulse on S, observing N+1 points (i,i) up to point (N,N);
 - (305) performing a pulse I (i.e. 180° pulse on I), which leads to (N,-N);
 - (310) waiting one time unit, leading to (N+1,-N+1);
 - (315) performing pulse IS (a 180° pulse on both I and S), leading to (-N-1,-N+1);
 - (320) observing points up to (N-2,N);
 - (325) performing a 180° pulse on I, leading to (N-2,-N);
 - (330) observing points up to (N+1,-N+3).
- 15. (Currently amended) The NMR spectroscopy method according to claim 1, wherein the pulse sequence spans a trajectory in a two-dimensional evolution time space (t_S, t_{IS}), said in performed on a hyperpolarized sample which comprises a first nuclear spin species (I) and a second nuclear spin species (S), the pulse sequence comprises the step of::
 - (300b) starting from the point (0,0), with an 90° pulse on S, observing N+1 points (i,i) up to point (N,N);
 - (305b) performing a pulse I (i.e. 180° pulse on I), which leads to (N,-N);
 - (310b) waiting one time unit, leading to (N+1,-N+1);

- (315b) performing a pulse IS (a-180° pulse on both I and S), leading to (-N-1,-N+1);
- (320b) observing points up to (N-2,N);
- (340b) performing a $\underline{180^{\circ}}$ pulse $\underline{\text{on }}$ S, which reverses both time signs and leads to (-N+2,-N);
- (345b) observing points up to (N,N-2);
- (350b) performing a 180° pulse on I leading to (N,-N+2);
- (355b) observing points up to (N+1,-N+3).
- 16. (New) A method of performing two-dimensional Nuclear Magnetic Resonance (NMR) spectroscopy on a hyperpolarized sample, which method comprises the steps of:
 - hyperpolarizing a sample which comprises a first nuclear species (I) and a second nuclear species (S), with the Hamiltonian $H = H_S + H_{IS} + H_{I}$ using DNP, wherein the NMR active nuclei receive hyperpolarization and transformation of the sample to a liquid state;
 - performing two-dimensional NMR spectroscopy on the sample and thereby producing at least two NMR spectra with the use of a sequence of rf-pulses, wherein pulse sequence comprises at least two rf-pulses on different nuclei, and wherein pulse sequence is adapted for a hyperpolarized sample in such a way that it uses a single scan, an efficient trajectory in a t_{S} - t_{IS} plane and produces a square array of observed points in a square portion of a two time space,;
 - analyzing the at least two NMR spectra in order to obtain a characterization of the sample.
- 17. (New) The NMR spectroscopy method according to claim 16, wherein the pulse sequence spans a trajectory in a two-dimensional evolution time space (t_S, t_{IS}), said pulse sequence comprises the step of:
 - (300) starting from the point (0,0), with an 90° pulse on S, observing N+1 points (i,i) up to point (N,N);
 - (305) performing a 180° pulse on I, which leads to (N,-N);
 - (310) waiting one time unit, leading to (N+1,-N+1);
 - (315) performing a 180° pulse on both I and S, leading to (-N-1,-N+1);

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- (320) observing points up to (N-2,N);
- (325) performing a 180° pulse on I, leading to (N-2,-N);
- (330) observing points up to (N+1,-N+3).
- 18. (New) The NMR spectroscopy method according to claim 16, wherein the pulse sequence spans a trajectory in a two-dimensional evolution time space (t_S, t_{IS}), said pulse sequence comprises the step of::
 - (300b) starting from the point (0,0), with an 90° pulse on S, observing N+1 points (i,i) up to point (N,N);
 - (305b) performing a 180° pulse on I, which leads to (N,-N);
 - (310b) waiting one time unit, leading to (N+1,-N+1);
 - (315b) performing a 180° pulse on both I and S, leading to (-N-1,-N+1);
 - (320b) observing points up to (N-2,N);
 - (340b) performing a 180° pulse on S, which reverses both time signs and leads to (-N+2,-N);
 - (345b) observing points up to (N,N-2);
 - (350b) performing a 180° pulse on I leading to (N,-N+2);
 - (355b) observing points up to (N+1,-N+3).